RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _08/444,791A_

Source: _1Fw/6_

Date Processed by STIC: _3/22/05_

ENTERED

RAW SEQUENCE LISTING   DATE: 03/22/2005
PATENT APPLICATION: US/08/444,791A   TIME: 11:53:10

Input Set : A:\40451C.txt
Output Set: N:\CRF4\03222005\H444791A.raw

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 3 <110> APPLICANT: Brockhaus, et al.
 5 <120> TITLE OF INVENTION: Human TNF Receptor
 7 <130> FILE REFERENCE: 01017/40451C
 9 <140> CURRENT APPLICATION NUMBER: US 08/444,791A
10 <141> CURRENT FILING DATE: 1995-05-19
12 <160> NUMBER OF SEQ ID NOS: 26
14 <170> SOFTWARE: PatentIn version 3.3
16 <210> SEQ ID NO: 1
17 <211> LENGTH: 2111
18 <212> TYPE: DNA
19 <213> ORGANISM: Homo sapiens
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26 ccagcactgc cgctgccaca ctgccctgag cccaaatggg ggagtgagag gccatagctg     180
28 tctggcatgg gcctctccac cgtgcctgac ctgctgctgc cgctggtgct cctggagctg     240
30 ttggtgggaa tatacccctc aggggttatt ggactggtcc ctcacctagg gacagggag     300
32 aagagagata gtgtgtgtcc ccaaggaaaa tatatccacc ctcaaaataa ttcgatttgc     360
34 tgtaccaagt gccacaaagg aacctacttg tacaatgact gtccaggccc ggggcaggat     420
36 acggactgca gggagtgtga gagcggctcc ttcaccgctt cagaaaacca cctcagacac     480
38 tgcctcagct gctccaaatg ccgaaaggaa atgggtcagg tggagatctc ttcttgcaca     540
40 gtggaccggg acaccgtgtg tggctgcagg aagaaccagt accggcatta ttggagtgaa     600
42 aacctttcc agtgcttcaa ttgcagcctc tgcctcaatg gaccgtgca cctctcctgc     660
44 caggagaaac agaacaccgt gtgcacctgc catgcaggtt tctttctaag agaaaacgag     720
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48 attgagaatg ttaagggcac tgaggactca ggcaccacag tgctgttgcc cctggtcatt     840
50 ttctttggtc tttgcctttt atccctcctc ttcattggtt taatgtatcg ctaccaacgg     900
52 tggaagtcca agctctactc cattgtttgt gggaaatcga cacctgaaaa agaggggag     960
54 cttgaaggaa ctactactaa gcccctggcc ccaaacccaa gcttcagtcc cactccaggc    1020
56 ttcacccca ccctgggctt cagtcccgtg cccagttcca ccttcacctc cagctccacc    1080
58 tatacccccg gtgactgtcc caactttgcg gctccccgca gagaggtggc accacctat    1140
60 cagggggctg accccatcct tgcgacagcc ctcgcctccg accccatccc caacccctt    1200
62 cagaagtggg aggacagcgc ccacaagcca cagagcctag acactgatga ccccgcgacg    1260
64 ctgtacgccg tggtggagaa cgtgcccccg ttgcgctgga ggaattcgt gcggcgccta    1320
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72 gaggcgcttt gcggccccgc cgccctcccg cccgcgccca gtcttctcag atgaggctgc    1560
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RAW SEQUENCE LISTING   DATE: 03/22/2005
PATENT APPLICATION: US/08/444,791A   TIME: 11:53:10

Input Set : A:\40451C.txt
Output Set: N:\CRF4\03222005\H444791A.raw

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86 gttttgtttt taaatcaatc atgttacact aatagaaact tggcactcct gtgccctctg    1980
88 cctggacaag cacatagcaa gctgaactgt cctaaggcag gggcgagcac ggaacaatgg    2040
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96 <211> LENGTH: 455
97 <212> TYPE: PRT
98 <213> ORGANISM: Homo sapiens
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107             20                  25                  30
110 His Leu Gly Asp Arg Glu Lys Arg Asp Ser Val Cys Pro Gln Gly Lys
111         35                  40                  45
114 Tyr Ile His Pro Gln Asn Asn Ser Ile Cys Cys Thr Lys Cys His Lys
115     50                  55                  60
118 Gly Thr Tyr Leu Tyr Asn Asp Cys Pro Gly Pro Gly Gln Asp Thr Asp
119 65                  70                  75                  80
122 Cys Arg Glu Cys Glu Ser Gly Ser Phe Thr Ala Ser Glu Asn His Leu
123                 85                  90                  95
126 Arg His Cys Leu Ser Cys Ser Lys Cys Arg Lys Glu Met Gly Gln Val
127             100                 105                 110
130 Glu Ile Ser Ser Cys Thr Val Asp Arg Asp Thr Val Cys Gly Cys Arg
131         115                 120                 125
134 Lys Asn Gln Tyr Arg His Tyr Trp Ser Glu Asn Leu Phe Gln Cys Phe
135     130                 135                 140
138 Asn Cys Ser Leu Cys Leu Asn Gly Thr Val His Leu Ser Cys Gln Glu
139 145                 150                 155                 160
142 Lys Gln Asn Thr Val Cys Thr Cys His Ala Gly Phe Phe Leu Arg Glu
143             165                 170                 175
146 Asn Glu Cys Val Ser Cys Ser Asn Cys Lys Lys Ser Leu Glu Cys Thr
147             180                 185                 190
150 Lys Leu Cys Leu Pro Gln Ile Glu Asn Val Lys Gly Thr Glu Asp Ser
151         195                 200                 205
154 Gly Thr Thr Val Leu Leu Pro Leu Val Ile Phe Phe Gly Leu Cys Leu
155     210                 215                 220
158 Leu Ser Leu Leu Phe Ile Gly Leu Met Tyr Arg Tyr Gln Arg Trp Lys
159 225                 230                 235                 240
162 Ser Lys Leu Tyr Ser Ile Val Cys Gly Lys Ser Thr Pro Glu Lys Glu
163             245                 250                 255
166 Gly Glu Leu Glu Gly Thr Thr Thr Lys Pro Leu Ala Pro Asn Pro Ser
167             260                 265                 270
170 Phe Ser Pro Thr Pro Gly Phe Thr Pro Thr Leu Gly Phe Ser Pro Val
171         275                 280                 285
174 Pro Ser Ser Thr Phe Thr Ser Ser Ser Thr Tyr Thr Pro Gly Asp Cys
175     290                 295                 300
178 Pro Asn Phe Ala Ala Pro Arg Arg Glu Val Ala Pro Pro Tyr Gln Gly
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187             340                   345                   350
190 Thr Asp Asp Pro Ala Thr Leu Tyr Ala Val Val Glu Asn Val Pro Pro
191         355                   360                   365
194 Leu Arg Trp Lys Glu Phe Val Arg Arg Leu Gly Leu Ser Asp His Glu
195     370                   375                   380
198 Ile Asp Arg Leu Glu Leu Gln Asn Gly Arg Cys Leu Arg Glu Ala Gln
199 385                   390                   395                   400
202 Tyr Ser Met Leu Ala Thr Trp Arg Arg Arg Thr Pro Arg Arg Glu Ala
203             405                   410                   415
206 Thr Leu Glu Leu Leu Gly Arg Val Leu Arg Asp Met Asp Leu Leu Gly
207             420                   425                   430
210 Cys Leu Glu Asp Ile Glu Glu Ala Leu Cys Gly Pro Ala Ala Leu Pro
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214 Pro Ala Pro Ser Leu Leu Arg
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219 <211> LENGTH: 2339
220 <212> TYPE: DNA
221 <213> ORGANISM: Homo sapiens
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228 actcgggaac agaaccgcat ctgcacctgc aggcccggct ggtactgcgc gctgagcaag     180
230 caggaggggt gccggctgtg cgcgccgctg ccgaagtgcc gcccgggctt cggcgtggcc     240
232 agaccaggaa ctgaaacatc agacgtggtg tgcaagccct gtgccccggg gacgttctcc     300
234 aacacgactt catccacgga tatttgcagg ccccaccaga tctgtaacgt ggtggccatc     360
236 cctgggaatg caagcaggga tgcagtctgc acgtccacgt cccccacccg gagtatggcc     420
238 ccaggggcag tacacttacc ccagccagtg tccacacgat cccaacacac gcagccaagt     480
240 ccagaaccca gcactgctcc aagcacctcc ttcctgctcc caatgggccc cagcccccca     540
242 gctgaaggga gcactggcga cttcgctctt ccagttggac tgattgtggg tgtgacagcc     600
244 ttgggtctac taataatagg agtggtgaac tgtgtcatca tgacccaggt gaaaaagaag     660
246 cccttgtgcc tgcagagaga agccaaggtg cctcacttgc ctgccgataa ggcccggggt     720
248 acacagggcc ccgagcagca gcacctgctg atcacagcgc cgagctccag cagcagctcc     780
250 ctggagagct cggccagtgc gttggacaga agggcgccca ctcggaacca gccacaggca     840
252 ccaggcgtgg aggccagtgg ggccggggag gcccgggcca gcaccgggag ctcagcagat     900
254 tcttcccctg gtggccatgg acccaggtc aatgtcacct gcatcgtgaa cgtctgtagc     960
256 agctctgacc acagctcaca gtgctcctcc caagccagct ccacaatggg agacacagat    1020
258 tccagcccct cggagtcccc gaaggacgag caggtcccct tctccaagga ggaatgtgcc    1080
260 tttcggtcac agctggagac gccagagacc ctgctgggga gcaccgaaga gaagcccctg    1140
262 cccccttggag tgcctgatgc tgggatgaag cccagttaac caggccggtg tgggctgtgt    1200
264 cgtagccaag gtggctgagc cctggcagga tgaccctgcg aaggggccct ggtccttcca    1260
266 ggcccccacc actaggactc tgaggctctt tctgggccaa gttcctctag tgccctccac    1320
268 agccgcagcc tccctctgac ctgcaggcca agagcagagg cagcgagttg tggaaagcct    1380
270 ctgctgccat ggcgtgtccc tctcggaagg ctggctgggc atggacgttc ggggcatgct    1440
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274 ctggagccct tgggtttttt gtttgtttgt ttgtttgttt gtttgtttct cccctgggc   1560
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278 agaggaggga tgctgcctga gtcacccatg aagacaggac agtgcttcag cctgaggctg   1680
280 agactgcggg atggtcctgg ggctctgtgc agggaggagg tggcagccct gtagggaacg   1740
282 gggtccttca agttagctca ggaggcttgg aaagcatcac ctcaggccag gtgcagtggc   1800
284 tcacgcctat gatcccagca ctttgggagg ctgaggcggg tggatcacct gaggttagga   1860
286 gttcgagacc agcctggcca acatggtaaa accccatctc tactaaaaat acagaaatta   1920
288 gccgggcgtg gtggcgggca cctatagtcc cagctactca gaagcctgag gctgggaaat   1980
290 cgtttgaacc cgggaagcgg aggttgcagg gagccgagat cacgccactg cactccagcc   2040
292 tgggcgacag agcgagagtc tgtctcaaaa gaaaaaaaaa aagcaccgcc tccaaatgct   2100
294 aacttgtcct tttgtaccat ggtgtgaaag tcagatgccc agagggccca ggcaggccac   2160
296 catattcagt gctgtggcct gggcaagata acgcacttct aactagaaat ctgccaattt   2220
298 tttaaaaaag taagtaccac tcaggccaac aagccaacga caaagccaaa ctctgccagc   2280
300 cacatccaac cccccacctg ccatttgcac cctccgcctt cactccggtg tgcctgcag   2339
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304 <211> LENGTH: 392
305 <212> TYPE: PRT
306 <213> ORGANISM: Homo sapiens
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314 Trp Asn Trp Val Pro Glu Cys Leu Ser Cys Gly Ser Arg Cys Ser Ser
315             20                  25                  30
318 Asp Gln Val Glu Thr Gln Ala Cys Thr Arg Glu Gln Asn Arg Ile Cys
319         35                  40                  45
322 Thr Cys Arg Pro Gly Trp Tyr Cys Ala Leu Ser Lys Gln Glu Gly Cys
323     50                  55                  60
326 Arg Leu Cys Ala Pro Leu Pro Lys Cys Arg Pro Gly Phe Gly Val Ala
327 65                  70                  75                  80
330 Arg Pro Gly Thr Glu Thr Ser Asp Val Val Cys Lys Pro Cys Ala Pro
331             85                  90                  95
334 Gly Thr Phe Ser Asn Thr Thr Ser Ser Thr Asp Ile Cys Arg Pro His
335             100                 105                 110
338 Gln Ile Cys Asn Val Val Ala Ile Pro Gly Asn Ala Ser Arg Asp Ala
339         115                 120                 125
342 Val Cys Thr Ser Thr Ser Pro Thr Arg Ser Met Ala Pro Gly Ala Val
343     130                 135                 140
346 His Leu Pro Gln Pro Val Ser Thr Arg Ser Gln His Thr Gln Pro Ser
347 145                 150                 155                 160
350 Pro Glu Pro Ser Thr Ala Pro Ser Thr Ser Phe Leu Leu Pro Met Gly
351             165                 170                 175
354 Pro Ser Pro Pro Ala Glu Gly Ser Thr Gly Asp Phe Ala Leu Pro Val
355             180                 185                 190
358 Gly Leu Ile Val Gly Val Thr Ala Leu Gly Leu Leu Ile Ile Gly Val
359         195                 200                 205
362 Val Asn Cys Val Ile Met Thr Gln Val Lys Lys Lys Pro Leu Cys Leu
363     210                 215                 220
366 Gln Arg Glu Ala Lys Val Pro His Leu Pro Ala Asp Lys Ala Arg Gly
367 225                 230                 235                 240
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370 Thr Gln Gly Pro Glu Gln Gln His Leu Leu Ile Thr Ala Pro Ser Ser
371             245             250             255
374 Ser Ser Ser Ser Leu Glu Ser Ser Ala Ser Ala Leu Asp Arg Arg Ala
375             260             265             270
378 Pro Thr Arg Asn Gln Pro Gln Ala Pro Gly Val Glu Ala Ser Gly Ala
379         275             280             285
382 Gly Glu Ala Arg Ala Ser Thr Gly Ser Ser Ala Asp Ser Ser Pro Gly
383     290             295             300
386 Gly His Gly Thr Gln Val Asn Val Thr Cys Ile Val Asn Val Cys Ser
387 305             310             315                 320
390 Ser Ser Asp His Ser Ser Gln Cys Ser Ser Gln Ala Ser Ser Thr Met
391             325             330             335
394 Gly Asp Thr Asp Ser Ser Pro Ser Glu Ser Pro Lys Asp Glu Gln Val
395             340             345             350
398 Pro Phe Ser Lys Glu Glu Cys Ala Phe Arg Ser Gln Leu Glu Thr Pro
399         355             360             365
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407 385                 390
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411 <211> LENGTH: 28
412 <212> TYPE: PRT
413 <213> ORGANISM: Artificial sequence
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416 <223> OTHER INFORMATION: Synthetic peptide
419 <220> FEATURE:
420 <221> NAME/KEY: misc_feature
421 <222> LOCATION: (25)..(25)
422 <223> OTHER INFORMATION: Xaa = unknown amino acid
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431             20              25
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434 <211> LENGTH: 15
435 <212> TYPE: PRT
436 <213> ORGANISM: Artificial sequence
438 <220> FEATURE:
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441 <400> SEQUENCE: 6
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447 <210> SEQ ID NO: 7
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449 <212> TYPE: PRT
450 <213> ORGANISM: Artificial sequence
452 <220> FEATURE:
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 03/22/2005
PATENT APPLICATION: US/08/444,791A   TIME: 11:53:11

Input Set : **A:\40451C.txt**
Output Set: **N:\CRF4\03222005\H444791A.raw**

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; Xaa Pos. 25
Seq#:10; Xaa Pos. 8
Seq#:11; Xaa Pos. 2
Seq#:14; Xaa Pos. 9,10,13

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L:430 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:16
L:509 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:533 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
L:586 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0
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